CSEP 524: Parallel Computation (week 8)

Brad Chamberlain

Tuesdays 6:30 – 9:20

MGH 231



Partitioned Global Address Space (PGAS) Programming Models



Partitioned Global Address Space Languages

(Or perhaps: partitioned global namespace languages)

- abstract concept:
 - support a shared namespace on distributed memory
 - permit any parallel task to access any lexically visible variable
 - doesn't matter if it's local or remote
 - establish a strong sense of ownership
 - every variable has a well-defined location
 - local variables are cheaper to access than remote ones

shared name-/address space					
private	private	private	private	private	
space 0	space 1	space 2	space 3	space 4	

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	partitioned s	shared name-/a	ddress space	
private	private	private	private	private
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Co-Array Fortran (CAF)

CAF: The first of our "traditional" PGAS languages

- developed ~1994
- adopted into the 2008 Fortran standard

Motivating Philosophy: "What is the smallest change required to convert Fortran 95 into a robust parallel language?"

originally referred to as F-- to emphasize "smallest change"

Quick Fortran Review/Intro

Traditional variables in Fortran:

```
! declares an integer, i

real x ! declares a float, x

real a(20) ! declares a 20-element array

real b(N,N) ! declares an N x N array
```

Array accesses are written with parenthesis:

```
a(1) = x ! Fortran uses 1-based indexing by default b(1,1) = 2*x b(2,:) = 3*x ! assign 3*x to the second row of b ! (':' is like '..' in Chapel)
```

CAF is **SPMD**

- SPMD programming/execution model
 - similar to MPI* in this regard
 - program copies are referred to as 'images'
- Use intrinsic functions to query the basics:

```
integer :: p, me
p = num_images() ! returns number of processes
me = this_image() ! returns value in 1..num_images()
```

• Barrier sync:

```
sync_all() ! wait for all processes/images
```

*= typical uses of it, anyway



Main CAF Concept: Co-Dimensions

Co-Dimension: an array dimension that refers to the space of CAF *images* (processes)

- defined using square brackets
 - (distinguishes it syntactically from a traditional dimension)

Main CAF Concept: Co-Dimensions

Co-array variables in Fortran:

```
integer i[*] ! declares an integer, i, per image
real x[*] ! declares a float, x, per image
real a(20)[*] ! declares a 20-element array per image
real b(N,N)[*] ! declares an N x N array per image
```

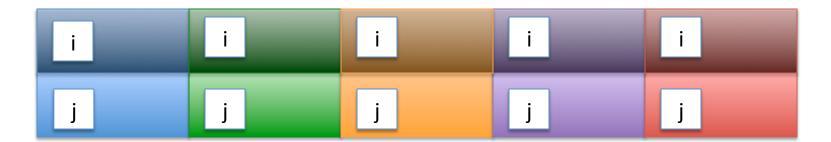
Main CAF Concept: Co-Dimensions

Co-array variables in Fortran:

```
integer i[*] ! declares an integer, i, per image
```

 Of course, traditional variables also result in a copy per image (it's SPMD after all), but private to that image

```
integer j ! declares a private integer, j, per image
```





Using Co-Arrays

```
integer i[*]
real x[*]
```

Refer to other images' values via co-array indexing:

```
if (me == 2) then
  nextX = x[me+1] ! read neighbor's value of x
  i[1] = i ! copy my value of 'i' into image 1's
endif
```

Co-array indexing/square brackets ⇒ communication

Stylized Collective Communications in CAF

Given declarations:

```
real x[*] real y real a(num_images())
```

Broadcast:

$$x[:] = y$$

Reduction:

$$y = MINVAL(x[:])$$

Gather:

$$a(:) = x[:]$$

Scatter:

$$x[:] = a(:)$$

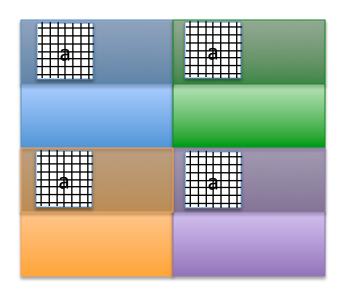


- When things divide evenly, you're pretty happy:
 - e.g., 1000 x 1000 array on a 2 x 2 processor grid:

```
real a (500,500) [2,2]
```

– or, adding in additional space for stencil ghost cells:

```
real a(0:501, 0:501)[2,2]
```





- When things divide evenly, you're pretty happy:
 - e.g., 1000 x 1000 array on a 2 x 2 processor grid:

```
real a (500,500) [2,2]
```

— or, adding in additional space for stencil ghost cells:

```
real a (0:501, 0:501) [2,2]
```

Stencil-style boundary value communication idioms:

```
! compute myrow, mycol, numrows, numcols
  if (myrow .ne. 1) then
    a(0,:) = a(500,:) [myrow-1, mycol]
  endif
  if (myrow .ne. numrows) then
    a(501,:) = a(1,:) [myrow+1, mycol]
  endi f
  ! etc.
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```



- When they don't, more work is required...
 - e.g., 1000 x 1000 array on a 2 x 3 processor grid:

```
real a (500, 334) [2, 3] ! allocate ceil(n/p) everywhere ...and then the images have to do bookkeeping to keep track of which image(s) own 334 items and which own 333
```

- details start to resemble the 9-point MPI code from HW
 - e.g., global-to-local and local-to-global index transformations
 - also, due to PGAS model, need to know more about neighbors
 - MPI: "I'll send you my high column which has index 333!"; "I'll recv it!"
 - CAF: "I'm going to access your high column" ⇒ "I must know its index"
 - (of course, some of this applies when things divide evenly as well...)



CAF Summary

- Program in SPMD style
- Communicate via variables with co-dimensions
 - a copy per program image
 - refer to other images' copies via square bracket subscripts
 - take advantage of good multidimensional array support
 - multidimensional views of process grid
 - multidimensional views of local data
 - syntactic support for slicing (:)
- Other stuff too, but this gives you the main idea
- Adopted into Fortran 2008 standard
 - see also http://www.co-array.org



CAF 2.0 (Rice University)

Motivation: Respond to a lack of richness in CAF

- difficult to have sets of images doing distinct things (teams)
- no support for pointer-based data structures
- poor support for collectives

For more information:

http://caf.rice.edu



UPC: Unified Parallel C

UPC: Our second "traditional" PGAS language

- developed ~1999
- "unified" in the sense that it combined 3 distinct parallel Cs:
 - AC, Split-C, Parallel C Preprocessor
- though a sibling to CAF, philosophically quite different

Motivating Philosophy:

- extend C concepts logically to support SPMD execution
 - 1D arrays
 - for loops
 - pointers (and pointer/array equivalence)



UPC is also **SPMD**

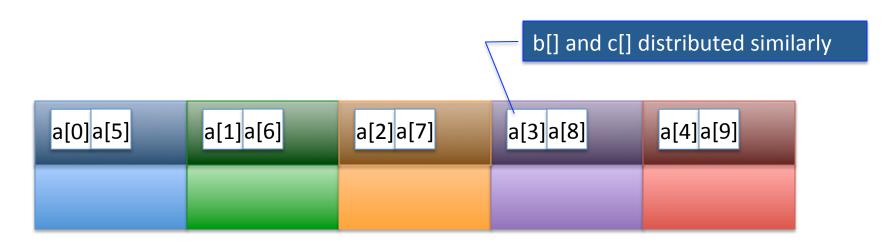
- SPMD programming/execution model
 - program copies are referred to as 'threads'
- Built-in constants provide the basics:

```
int p, me;
p = THREADS;  // returns number of processes
me = MYTHREAD;  // returns a value in 0..THREADS-1
```

Barrier synch statement:

- Arrays declared with the 'shared' keyword are distributed within the shared space
 - uses a cyclic distribution by default

```
#define N 10
shared float a[N], b[N], c[N];
```

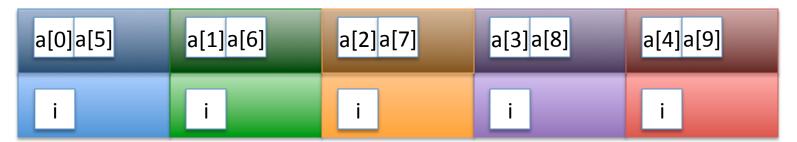




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- Arrays declared with the 'shared' keyword are distributed within the shared space
 - uses a cyclic distribution by default

```
#define N 10
shared float a[N], b[N], c[N];
for (int i=0; i<N; i++) { // dumb loop: O(N)
  if (i%THREADS == MYTHREAD) {
    c[i] = a[i] + alpha * b[i];
}</pre>
```





- Arrays declared with the 'shared' keyword are distributed within the shared space
 - uses a cyclic distribution by default

```
#define N 10
shared float a[N], b[N], c[N];
// smarter loop: O(N/THREADS)

for (int i=MYTHREAD; i<N; i+=THREADS) {
   c[i] = a[i] + alpha * b[i];
}</pre>
```

a[0]a[5]	a[1] <mark>a[6]</mark>	a[2]a[7]	a[3]a[8]	a[4]a[9]
i	i	i	i	i



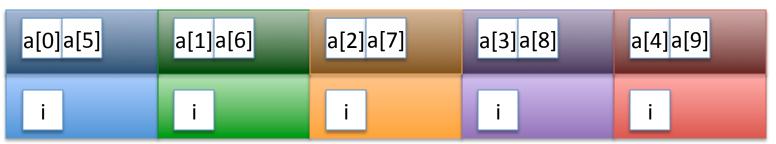
 Arrays declared with the 'shared' keyword are distributed within the shared space

```
- uses a cyclic distribution by default
  #define N 10
  shared float a[N], b[N], c[N];

// "global-view"equivalent to upc_forall (int i=0; i<N; i++; i) {
  c[i] = a[i] + alpha * b[i];
}</pre>
Affinity field: Which thread should execute this iteration?
(if int, %THREADS to get ID)

the previous

i) {
  c[i] = a[i] + alpha * b[i];
}
```





- Arrays declared with the 'shared' keyword are distributed within the shared space
 - can specify a block-cyclic distribution as well

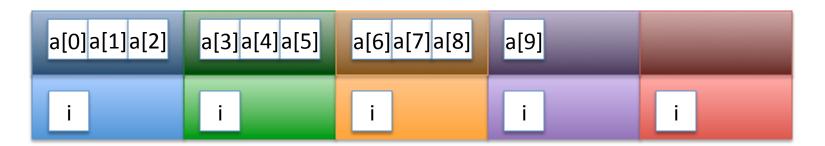
```
#define N 10
shared [2] float a[N], b[N], c[N];
upc_forall (int i=0; i<N; i++; &c[i]) {
   c[i] = a[i] + alpha * b[i];
}
Affinity field: Which thread
should execute this iteration?
(if ptr-to-shared, owner does)</pre>
```

a[0]a[1]	a[2] <mark>a[3]</mark>	a[4]a[5]	a[6] <mark>a[7]</mark>	a[8]a[9]
i	i	i	i	i



- Arrays declared with the 'shared' keyword are distributed within the shared space
 - can specify a block-cyclic distribution as well

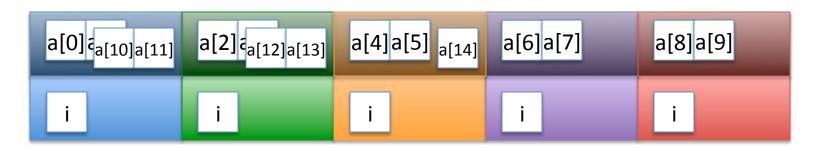
```
#define N 10
shared [3] float a[N], b[N], c[N];
upc_forall (int i=0; i<N; i++; &c[i]) {
   c[i] = a[i] + alpha * b[i];
}</pre>
```





- Arrays declared with the 'shared' keyword are distributed within the shared space
 - can specify a block-cyclic distribution as well

```
#define N 15
shared [2] float a[N], b[N], c[N];
upc_forall (int i=0; i<N; i++; &c[i]) {
   c[i] = a[i] + alpha * b[i];
}</pre>
```





Scalars in UPC

 Somewhat confusingly (to me anyway*), shared scalars in UPC result in a single copy on thread 0

```
int i;
shared int j;
```

^{* =} because it seems contrary to SPMD programming

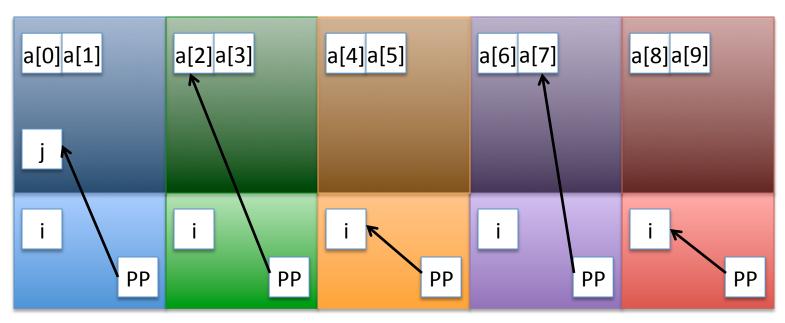




Pointers in UPC

 UPC Pointers may be private/shared and may point to private/shared

int* PP; // private pointer to local data

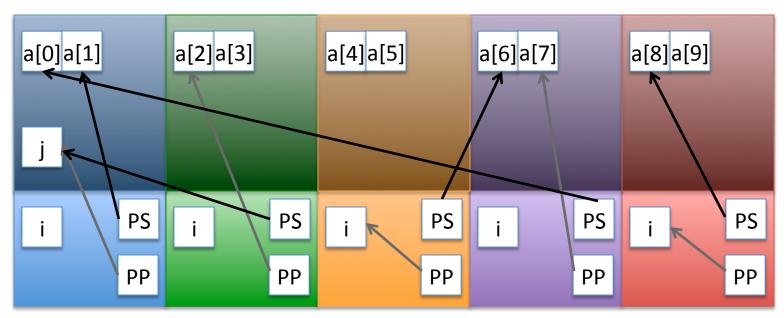




Pointers in UPC

 UPC Pointers may be private/shared and may point to private/shared

```
int* PP; // private pointer to local data
shared int* PS; // private pointer to shared data
```

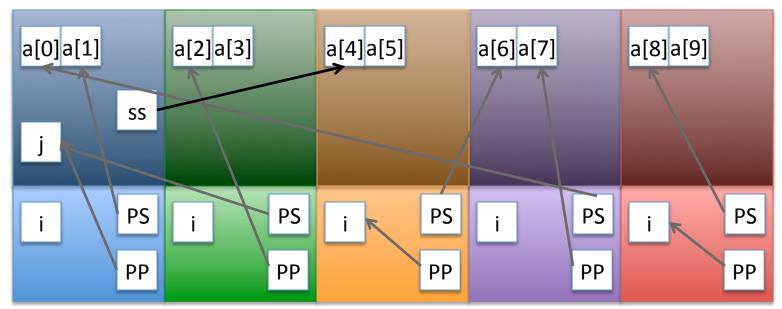




Pointers in UPC

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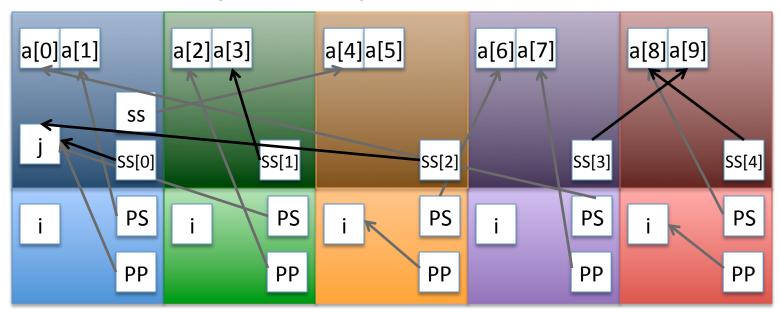
```
int* PP; // private pointer to local data
shared int* PS; // private pointer to shared data
shared int* shared ss; // shared pointer to shared data
```





Arrays of Pointers in UPC

- Of course, one can also create arrays of pointers
 // array of shared pointer to shared data
 shared int* shared SS[THREADS];
- As you can imagine, one UPC's strengths is its ability to create fairly arbitrary distributed data structures

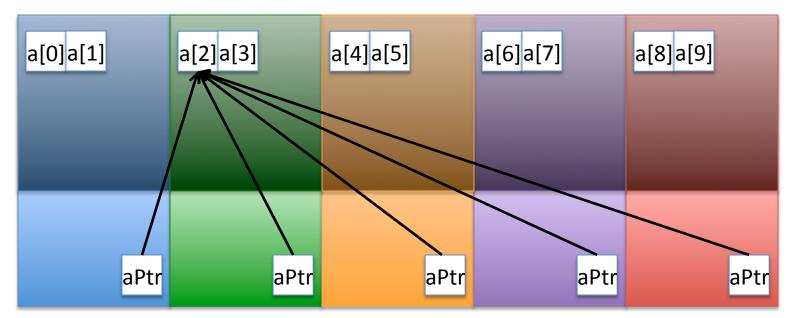




Array/Pointer Equivalence in UPC

As in C, pointers can be walked through memory

```
shared [2] float a[N];
shared [2] float* aPtr = &(a[2]);
```

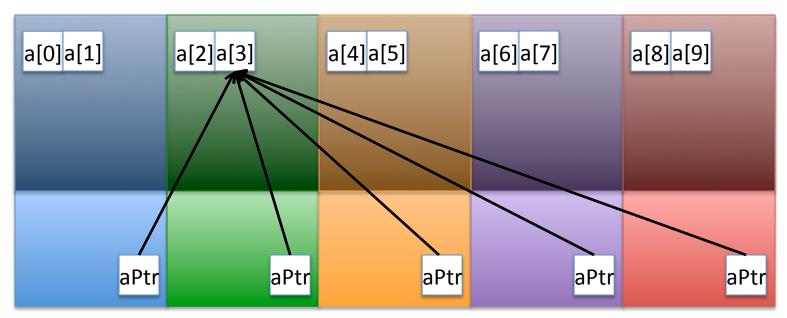




Array/Pointer Equivalence in UPC

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shared [2] float* aPtr = &(a[2]);
aPtr++;
```

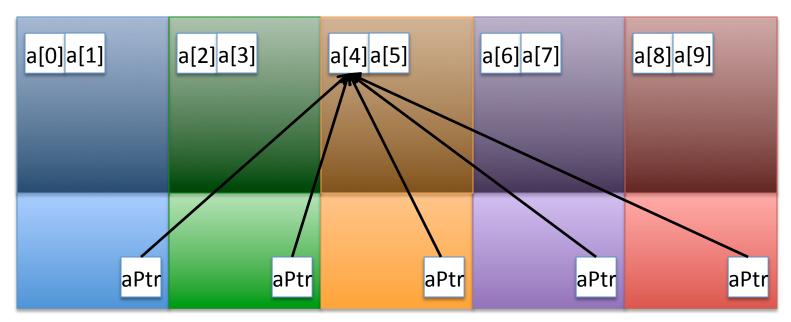




Array/Pointer Equivalence in UPC

As in C, pointers can be walked through memory

```
shared [2] float a[N];
shared [2] float* aPtr = &(a[2]);
aPtr++;
aPtr++;
```





How are UPC Pointers Implemented?

Local pointers to local: just an address, as always

Pointers to shared: 3 parts

- thread ID
- base address of block within the thread
- phase/offset within the block (0..blocksize-1)
- UPC supports a number of utility functions that permit you to query this information from pointers
- Casting between pointer types is permitted
 - but can be dangerous (as in C) and/or lossy



UPC: Local-view or Global-view?

Global arrays and pointers: global-view

upc_forall loops: global-view

Shared scalars: global-view-ish (but constrained)

Private scalars: local-view

SPMD model: local-view

⇒ a bit of both

Other Features in UPC

- Collectives Library
- Memory Consistency Model
 - among the first/foremost memory models in HPC
 - ability to move between strict and relaxed models
 - fence operations
- Dynamic Memory Management
- Locks
- Parallel I/O
- ...



Titanium: Java-based PGAS language

Titanium: The third traditional PGAS language

- And in my opinion, the most promising in terms of features
- Based on Java, though loosely at times

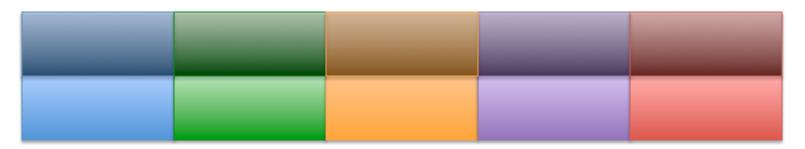
- Unfortunately didn't catch on as well
 - in part because Java not dominant in HPC
 - in part because of "superset of subset" problem
 - it's like Java except for when it's completely different
- Last I heard, "not quite dead yet"

PGAS: What's in a Name?

		memory model	programming model	execution model	data structures	communication	
	MPI	distributed memory		executables) in practice)	manually fragmented	APIs	
	OpenMP	shared memory	global-view parallelism	shared memory multithreaded	shared memory arrays	N/A	
es	CAF				co-arrays	co-array refs	
PGAS Languages	UPC	PGAS	Single Program, Multiple Data (SPMD)		1D block-cyc arrays/ distributed pointers	implicit	
Lar	Titanium				class-based arrays/ distributed pointers	method-based	
	Chapel	PGAS	global-view parallelism	distributed memory multithreaded	global-view distributed arrays	implicit	



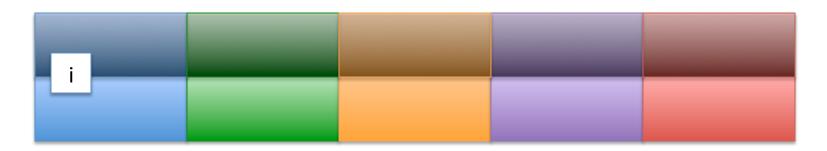
- Chapel differs from UPC/CAF since it's not SPMD
 - ⇒"global name-/address space" comes from lexical scoping
 - rather than: "We're all running the same program, so we must all have a variable named x"
 - as in traditional languages, each declaration yields one variable
 - stored on locale where task executes, not everywhere/thread 0
 - ⇒ user-level concept of locality is central to language
 - parallelism and locality are two distinct things
 - shouldn't think in terms of "that other copy of the program"





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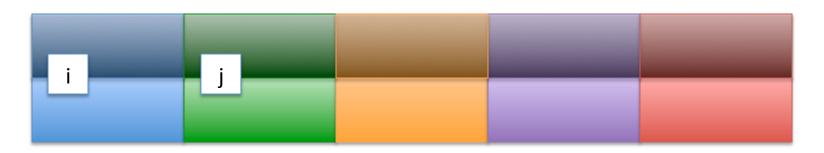
var i: int;





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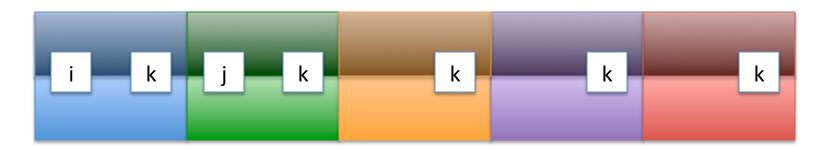
```
var i: int;
on Locales[1] {
  var j: int;
```





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```
var i: int;
on Locales[1] {
  var j: int;
  coforall loc in Locales {
    on loc {
     var k: int;
    }
}
```





Chapel and PGAS: Public vs. Private

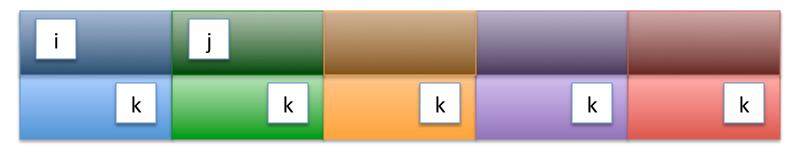
- How public a variable is depends only on scoping
 - who can see it?
 - who actually bothers to refer to it?

```
var i: int;
on Locales[1] {
  var j: int;
  coforall loc in Locales {
    on loc {
      var k = i + j;
          k
                      k
                                 k
                                             k
```



Chapel and PGAS: Public vs. Private

- How public a variable is depends only on scoping
 - who can see it?
 - who actually bothers to refer to it?
- Chapel represents variables that are referred to nonlocally using wide pointers
 - locale ID + local address
 - note: no need for phase/offset as in UPC
 - because no block-cyclic pointer math required





Single-Sided Communication



But First: Two-Sided Communication

two-sided communication: What we did in MPI

- one process sends a message
- another process receives
- both sides necessary for data to be transferred
 - else, deadlock

Implementing PGAS Languages: 1-sided comm.

single-sided (one-sided) communication: the backbone of most PGAS language implementations

primitive operations:

- get(): reads from a remote process's address space
- put(): writes to a remote process's address space
- No matching operation required!

Prototypical 1-sided comm. routines

(Many implementations will also support variations for strided puts/gets, multidimensional puts/gets, gather/scatter puts/gets)

Why does PGAS need/want 1-sided comm?

- Communication is expressed via naming variables that happen to live on another process
 - generally, one process will have no idea what other is doing
 - even in SPMD programming models
 - control flow may take different paths
 - local/private variables are likely to have different values
 - as a result, I can't guess what data of mine you might need
 - so I can't call the matching sends/recvs to fulfill your requests



Summary of 1-sided comm.

Characteristics:

- notably, the text of the remote program need do nothing
- in effect, implements load/store for non-trivial data sizes over distributed memory
- interestingly, has not become an end-user model like MPI
- key supporting network technology to work well: RDMA
 - Remote Direct Memory Access

• Benefits:

- results in fewer copies/buffers within the SW stack (often 0)
- separates data transfer from synchronization of processes
- with RDMA, doesn't require remote CPU to be involved



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Summary of 1-sided comm.

Drawbacks:

- if network has no RDMA support, performance can suffer
 - e.g., may require devoting a thread to handling incoming requests
 - (in particular, 1-sided comm. can be implemented using MPI)
- re-opens door to memory consistency issues

1-Sided Communication Implementations

SHMEM/OpenSHMEM (Cray/community)

the first (? major, anyway) single-sided comm. interface

GASNet (Berkeley)

(what Chapel uses by default)

ARMCI (PNNL)

GASPI (Germany)

MPI-3

as mentioned last week, part of newest feature set

Chapel's Extra Communication Requirement

In addition to puts/gets Chapel needs active messages

- "run this code over there with these arguments"
- can think of as a style of 1-sided communication
- active ⇒ control is transferred, not just data

Used to implement on-clauses

```
var i: int;
on Locales[1] {
    ... // send an active message to execute this code
}
```



Conceptual active message interface

Active Message Support?

SHMEM/OpenSHMEM (Cray/community)

GASNet (Berkeley)

ARMCI (PNNL)

GASPI (Germany)

MPI-3



Smith-Waterman Algorithm for Sequence Alignment



Goal: Determine the similarities/differences between two protein sequences/nucleotides.

e.g., ACACACTA and AGCACACA*

Basis of Computation: Defined via a recursive formula:

$$H(i,0) = 0$$

 $H(0,j) = 0$
 $H(i,j) = f(H(i-1, j-1), H(i-1, j), H(i, j-1))$

Caveat: This is a classic, rather than cutting-edge sequence alignment algorithm, but it illustrates an important parallel paradiagm: wavefront computation



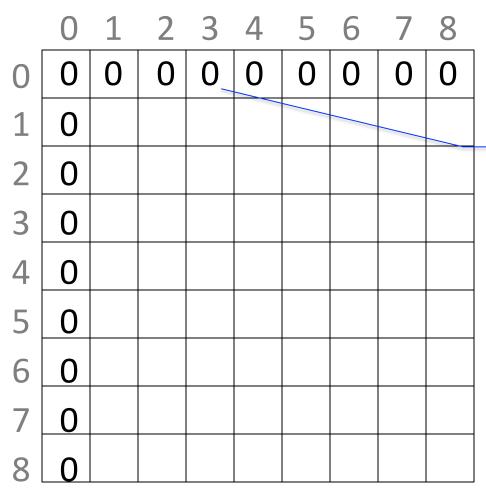
Naïve Task-Parallel Approach:

```
proc computeH(i, j) {
  if (i==0 | | i == 0) then
    return 0;
  else
    var h1, h2, h3: int;
    begin h1 = computeH(i-1, j-1);
    begin h2 = computeH(i-1, j);
    begin h3 = computeH(i, j-1);
    return f (h1, h2, h3);
```

Note: Recomputes most subexpressions redundantly

This is a case for dynamic programming!

Dynamic Programming Approach:

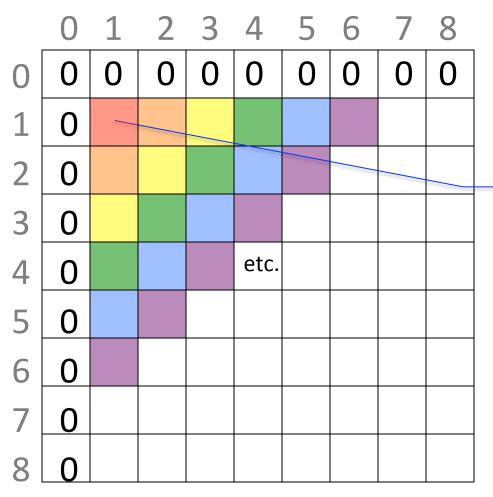


Step 1: Initialize boundaries to 0

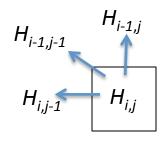


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Dynamic Programming Approach:



Step 2: Compute cells as we're able to





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Dynamic Programming Approach:

	0	1	2	3	4	5	6	7	8
0	0	0	0	0	0	0	0	0	0
1	0	2	1	2	1	2	1	0	2
2	0	1	1	1	1	1	1	0	1
3	0	0	3	2	3	2	3	2	1
4	0	2	2	5	4	5	4	3	4
5	0	1	4	4	7	6	7	6	5
6	0	2	3	6	6	9	8	7	8
7	0	1	4	5	8	8	11	10	9
8	0	2	3	6	7	10	10	10	12

Step 3: Follow trail of breadcrumbs back



Dynamic Programming Approach:

	0	1	2	3	4	5	6	7	8
0	0	0	0	0	0	0	0	0	0
1	0	2	1	2	1	2	1	0	2
2	0	1	1	1	1	1	1	0	1
3	0	0	3	2	3	2	3	2	1
4	0	2	2	5	4	5	4	3	4
5	0	1	4	4	7	6	7	6	5
6	0	2	3	6	6	9	8	7	8
7	0	1	4	5	8	8	11	10	9
8	0	2	3	6	7	10	10	10	12

Step 3: Follow trail of breadcrumbs back



Dynamic Programming Approach:

		Α	C	Α	C	Α	C	T	Α
	0	0	0	0	0	0	0	0	0
Α	0	2	1	2	1	2	1	0	2
G	0	1	1	1	1	1	1	0	1
C	0	0	3	2	3	2	3	2	1
Α	0	2	2	5	4	5	4	3	4
C	0	1	4	4	7	6	7	6	5
Α	0	2	3	6	6	9	8	7	8
С	0	1	4	5	8	8	11	10	9
Α	0	2	3	6	7	10	10	10	12

Step 4: Interpret the path against the original sequences

AGCACACAAA A-CACACTA

How could we do this in parallel?



Data-Parallel Approach:

```
Loop over upper diagonals serially
```

```
proc computeH(H: [0..n, 0..n] int) {
  for upperDiag in 1..n do
    forall diagPos in 0..#upperDiag {
      const (i,j) = [diagPos+1, upperDiag-diagPos];
      H[i,j] = f(H[i-1,j-1], H[i-1,j], H[i,j-1]);
    }
  for lowerDiag in 1..n-1 do
    forall diagPos in lowerDiag..n-1 by -1 {
      const (i,j) = [diagPos+1, lowerDiag+diagPos];
      H[i,j] = f(H[i-1,j-1], H[i-1,j], H[i,j-1]);
    }
}
```

Traverse each diagonal in parallel

Repeat for lower diagonals

Advantages:

Reasonably clean (if I got my indexing correct)

<u>Disadvantages:</u>

- Not so great in terms of cache use
- A bit fine-grained
 - max parallelism = N/P
- Not ideal for distributed memory



Naïve Data-Driven Task-Parallel Approach:

```
proc computeH(H: [0..n, 0..n] int) {
  const ProbSpace = H.domain.translate(1,1);
  var NeighborsDone: [ProbSpace] atomic int = 0;
  var Ready$: [ProbSpace] sync int;
 NeighborsDone[1, ..].add(1);
  NeighborsDone[.., 1].add(1);
  NeighborsDone[1, 1].add(1);
  Ready$[1,1] = 1;
  coforall (i, j) in ProbSpace {
    const goNow = Ready$[i,j];
    H[i,j] = f(H[i-1,j-1], H[i-1,j], H[i,j-1]);
    const eastReady = NeighborsDone[i, j+1].fetchAdd(1);
    const seReady = NeighborsDone[i+1,j+1].fetchAdd(1);
    const southReady = NeighborsDone[i+1, j].fetchAdd(1);
    if (eastReady == 2) then Ready\{[i,j+1] = 1;
    if (seReady == 2) then Ready\{[i+1,j+1] = 1;
    if (southReady == 2) then Ready$[i+1,j] = 1;
```

Create domain describing shifted version off H's domain

Arrays to count how many of our 3 neighbors are done; and to signal when we can compute

Set up boundaries: north and west elements have a neighbor done; top-left is ready

Create a task per matrix element and have it block until ready

Compute our element

Increment our neighbors' counts

Signal our neighbors as ready if we're the third



dSEP 524: Parallel Computation



Comparison of Synchronization Types in Chapel

sync/single:

- Best for producer/consumer style synchronization
- Imply a memory fence w.r.t. other loads/stores
- Use single to express write-once values

atomic:

Best for uncoordinated accesses to shared state





Naïve Data-Driven Task-Parallel Approach:

```
proc computeH(H: [0..n, 0..n] int) {
  const ProbSpace = H.domain.translate(1,1);
  var NeighborsDone: [ProbSpace] atomic int = 0;
  var Ready$: [ProbSpace] sync int;
  NeighborsDone[1, ..].add(1);
                                                  Disadvantages:
  NeighborsDone[.., 1].add(1);

    Still not great in cache use

  NeighborsDone[1, 1].add(1);

    Uses n<sup>2</sup> tasks

  Ready\{[1,1] = 1;
                                                     Most spend most of their
  coforall (i, j) in ProbSpace {
                                                     time blocking
    const goNow = Ready$[i, j];
    H[i,j] = f(H[i-1,j-1], H[i-1,j], H[i,j-1]);
    const eastReady = NeighborsDone[i,j+1].fetchAdd(1);
    const seReady = NeighborsDone[i+1,j+1].fetchAdd(1);
    const southReady = NeighborsDone[i+1, j].fetchAdd(1);
    if (eastReady == 2) then Ready\{[i,j+1] = 1;
    if (seReady == 2) then Ready\{[i+1,j+1] = 1;
    if (southReady == 2) then Ready$[i+1,j] = 1;
```



Slightly Less Naïve Data-Driven Task-Parallel Approach:

```
proc computeH(H: [0..n, 0..n] int) {
  const ProbSpace = H.domain.translate(1,1);
  var NeighborsDone: [ProbSpace] atomic int = 0;
  NeighborsDone[1, ..].add(1);
                                          Rather than create the tasks a priori, fire
  NeighborsDone[.., 1].add(1);
                                          them off once we know they're legal
  NeighborsDone[1, 1].add(1);
  sync { computeHHelp(1,1); }
                                     sync to ensure they're all done before we go on
 proc computeHHelp(i,j) {
    H[i,j] = f(H[i-1,j-1], H[i-1,j], H[i,j-1]);
    const eastReady = NeighborsDone[i,j+1].fetchAdd(1);
    const seReady = Neighborspone[i+1,j+1].fetchAdd(1);
    const southReady = NeighborsDone[i+1, j].fetchAdd(1);
    if (eastReady == 2) then begin computeHHelp(i, j+1);
    if (seReady == 2) then begin computeHHelp(i+1, j+1);
    if (southReady == 2) then begin computeHHelp(i+1, j);
```



Slightly Less Naïve Data-Driven Task-Parallel Approach:

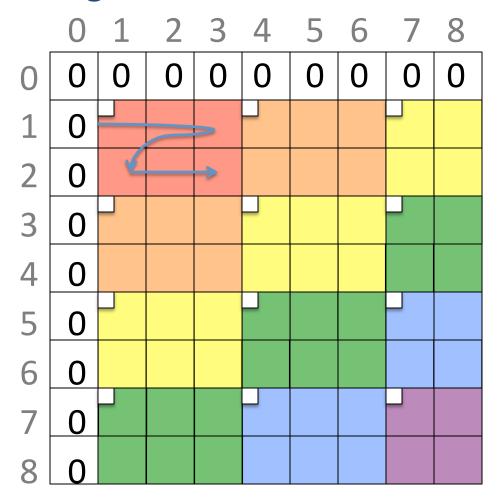
```
proc computeH(H: [0..n, 0..n] int) {
  const ProbSpace = H.domain.translate(1,1);
  var NeighborsDone: [ProbSpace] atomic int = 0;
  NeighborsDone[1, ..].add(1);
  NeighborsDone[.., 1].add(1);
                                                 Disadvantages:
  NeighborsDone[1, 1].add(1);

    Still uses a lot of tasks

  sync { computeHHelp(1,1); }
                                                    Each task is very fine-grained
 proc computeHHelp(i,j) {
    H[i,j] = f(H[i-1,j-1], H[i-1,j], H[i,j-1]);
    const eastReady = NeighborsDone[i,j+1].fetchAdd(1);
    const seReady = NeighborsDone[i+1, j+1].fetchAdd(1);
    const southReady = NeighborsDone[i+1, j].fetchAdd(1);
    if (eastReady == 2) then begin computeHHelp(i, j+1);
    if (seReady == 2) then begin computeHHelp(i+1,j+1);
    if (southReady == 2) then begin computeHHelp(i+1, j);
```



Coarsening the Parallelism:





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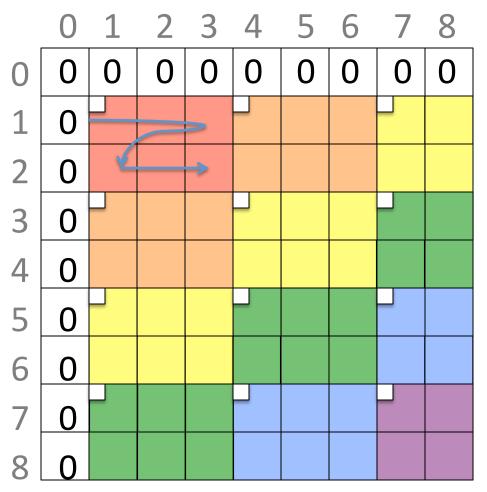
Stride indices to get to next chunk

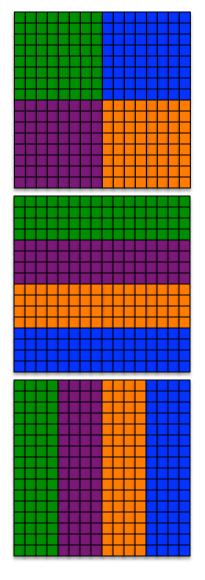
Blocked Data-Driven Task-Parallel Approach:

```
proc computeH(H: [0..n, 0..n] int) {
  const ProbSpace = H.domain.translate(1,1) by (rowsPerChunk, colsPerChunk);
  var NeighborsDone: [ProbSpace] atomic int = 0;
 NeighborsDone[1, ..].add(1);
                                              Can now use strided array for atomics
 NeighborsDone[.., 1].add(1);
  NeighborsDone[1, 1].add(1);
  sync { computeHHelp({1..rowsPerChunk,1..colsPerChunk}); }
                                               Change helper to take a domain
 proc computeHHelp(inds) {
                                              describing the chunk to compute
    for (i, j) in H.domain[inds]\do
      H[i,j] = f(H[i-1,j-1], H[i \uparrow 1,j], H[i,j-1]);
    const (i, j) = inds.low;
                                              Compute over chunk serially
    const eastReady = NeighborsDone(i, j+colsPerChunk).fetchAdd(1);
    const seReady = NeighborsDone[i+rowsPerChunk, j+colsPerChunk].fetchAdd(1);
    const southReady = NeighborsDone[i+rowsPerChunk, j].fetchAdd(1);
    if (eastReady == 2) then begin computeHHelp(i,j+colsPerChunk);
    if (seReady == 2) then begin computeHHelp(i+rowsPerChunk,j+colsPerChunk);
    if (southReadv == 2) then begin computeHHelp(i+rowsPerChunk,j);
```



Now, what about distributed memory?

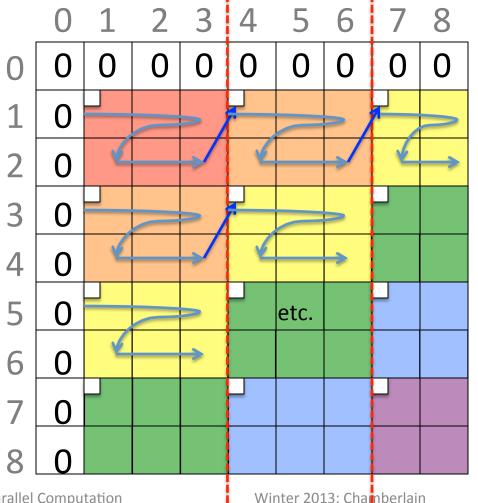






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Now, what about distributed memory?



Advantages:

- Good cache behavior: Nice fat blocks of data touchable in memory order
- Pipeline parallelism: Good utilization once pipeline is filled

Other notes:

- Communication pattern?
- Hybrid distributed + shared memory approach?



CSEP 524: Parallel Computation

Chapel Domain Maps

(switch to other slide deck)



From the Course Description...

styles of parallelism

- data-parallel
- task-parallel
- concurrency
- pipelined parallelism
- nested parallelism

abstract programming models

- shared memory
- Single Program, Multiple Data (SPMD)
- message passing
- Partitioned Global Address Space (PGAS)

architectural implications

- shared vs. distributed memory
- multicore processors and accelerators
- networks
- caches and memory

programming issues and hazards

- synchronization
- memory consistency
- race conditions
- deadlock and livelock

performance tuning

- scalability
- locality
- communication
- scalar concerns

programming languages and notations

- OpenMP
- MPI
- UPC
- Chapel
- CUDA/OpenCL/OpenACC (?)

algorithms and patterns

- reductions and scans
- stencils
- graph algorithms
- ..



Requests for next week?

- Amdahl's Law
- modern compute nodes: CPU+GPU, NUMA nodes
- software transactional memory
- ZPL/HPF: Grand failures of the 90's
- advanced Chapel concepts: user-defined arrays/foralls
- Dragonfly network
- open discussion questions
- more algorithms

